

## SEQUENCE LISTING

## (1) GENERAL INFORMATION:

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## (i) APPLICANT:

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- (A) NAME: Kanton Zuerich vertreten durch die Erziehungsdirektion
- (B) STREET: Walchetur
- (C) CITY: Zuerich
- (D) STATE: Zuerich
- (E) COUNTRY: Switzerland
- (F) POSTAL CODE (ZIP): CH-8090
- (G) TELEPHONE: +41-1-259 2388

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(ii) TITLE OF INVENTION: Immunological Detection of Prions

(iii) NUMBER OF SEQUENCES: 9

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## (iv) COMPUTER READABLE FORM:

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- (A) MEDIUM TYPE: Floppy disk
- (B) COMPUTER: IBM PC compatible
- (C) OPERATING SYSTEM: PC-DOS/MS-DOS
- (D) SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)

## (vi) PRIOR APPLICATION DATA:

- (A) APPLICATION NUMBER: EP 97102837.8
- (B) FILING DATE: 21-FEB-1997

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## (2) INFORMATION FOR SEQ ID NO: 1:

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## (i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 660 base pairs
- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: double
- (D) TOPOLOGY: linear

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## (ii) MOLECULE TYPE: DNA (genomic)

(iii) HYPOTHETICAL: NO

(iv) ANTI-SENSE: NO

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## (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Bos taurus
- (D) DEVELOPMENTAL STAGE: Adult

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## (vii) IMMEDIATE SOURCE:

- (B) CLONE: pbPrP3

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## (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:

ATGAAGAAC GACCAAAACC TGGAGGAGGA TGGAACACTG GGGGGAGCCG ATACCCAGGA 60  
CAGGGCAGTC CTGGAGGCAA CCCTTATCCA CCTCAGGGAG GGGGTGGCTG GGGTCAGCCC 120

CATGGAGGTG	GCTGGGGCCA	GCCTCATGGA	GGTGCCTGGG	CCCAGCCTCA	TGGACGTGGC	180	
TGGGCTCAGC	CCCATCGTGG	TGGCTGGGA	CAGCCACATG	GTGGTGGAGG	CTGGGGTCAA	240	
5	GGTGCCTACCC	ACGGTCAATG	GAACAAACCC	AGTAAGCCAA	AAACCAACAT	GAAGCATGTG	300
	GCAGGAGCTG	CTGCAGCTGG	ACCAGTGGTA	GGGGGCCCTTG	GTGGCTACAT	GCTGGGAAGT	360
10	GCCATGAGCA	GGCCTCTTAT	ACATTTGGC	AGTGACTATG	AGGACCGTTA	CTATCGTGAA	420
	AACATGCACC	GTTACCCCAA	CCAAGTGTAC	TACAGGCCAG	TGGATCAGTA	TAGTAACCAG	480
	AACAACTTG	TGCATGACTG	TGTCAACATC	ACAGTCAAGG	AACACACAGT	CACCACCACC	540
15	ACCAAGGGGG	AGAACTTCAC	CGAAACTGAC	ATCAAGATGA	TGGAGCGAGT	GGTGGAGCAA	600
	ATGTGCATTA	CCCAGTACCA	GAGAGAATCC	CAGGCTTATT	ACCAACGAGG	GGCAAGTTAA	660

## 20 (2) INFORMATION FOR SEQ ID NO: 2:

(i) SEQUENCE CHARACTERISTICS:

- (A) LENGTH: 219 amino acids
- (B) TYPE: amino acid
- 25 (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: protein

30 (iii) HYPOTHETICAL: YES

(iv) ANTI-SENSE: NO

35 (vi) ORIGINAL SOURCE:

- (A) ORGANISM: Bos taurus

(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 2:

40	Met Lys Lys Arg Pro Lys Pro Gly Gly Gly Trp Asn Thr Gly Gly Ser			
	1	5	10	15
45	Arg Tyr Pro Gly Gln Gly Ser Pro Gly Gly Asn Arg Tyr Pro Pro Gln			
	20	25	30	
50	Gly Gly Gly Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro			
	35	40	45	
55	His Gly Gly Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln Pro			
	50	55	60	
65	His Gly Gly Trp Gly Gln Pro His Gly Gly Gly Trp Gly Gln			
	65	70	75	80
75	Gly Gly Thr His Gly Gln Trp Asn Lys Pro Ser Lys Pro Lys Thr Asn			
	85	90	95	
85	Met Lys His Val Ala Gly Ala Ala Ala Gly Ala Val Val Gly Gly			
	100	105	110	

Leu Gly Gly Tyr Met Leu Gly Ser Ala Met Ser Arg Pro Leu Ile His  
 115 120 125  
 5 Phe Gly Ser Asp Tyr Glu Asp Arg Tyr Tyr Arg Glu Asn Met His Arg  
 130 135 140  
 Tyr Pro Asn Gln Val Tyr Tyr Arg Pro Val Asp Gln Tyr Ser Asn Gln  
 145 150 155 160  
 10 Asn Asn Phe Val His Asp Cys Val Asn Ile Thr Val Lys Glu His Thr  
 165 170 175  
 15 Val Thr Thr Thr Lys Gly Glu Asn Phe Thr Glu Thr Asp Ile Lys  
 180 185 190  
 Met Met Glu Arg Val Val Glu Gln Met Cys Ile Thr Gln Tyr Gln Arg  
 195 200 205  
 20 Glu Ser Gln Ala Tyr Tyr Gln Arg Gly Ala Ser  
 210 215

## (2) INFORMATION FOR SEQ ID NO: 3:

- 25 (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 33 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear  
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 (ii) MOLECULE TYPE: other nucleic acid  
 (A) DESCRIPTION: /desc = "oligonucleotide"  
 35 (iii) HYPOTHETICAL: NO  
 (iv) ANTI-SENSE: NO

40 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 3:  
 GGGATTCCA TATGAAGAAC CGACCAAAAC CTG

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## (2) INFORMATION FOR SEQ ID NO: 4:

- 45 (i) SEQUENCE CHARACTERISTICS:  
 (A) LENGTH: 31 base pairs  
 (B) TYPE: nucleic acid  
 (C) STRANDEDNESS: single  
 (D) TOPOLOGY: linear  
 50  
 (ii) MOLECULE TYPE: other nucleic acid  
 (A) DESCRIPTION: /desc = "oligonucleotide"  
 55 (iii) HYPOTHETICAL: NO  
 (iv) ANTI-SENSE: NO

5 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 4:

CGGGATCCTA TTAACTTGCC CCTCGTTGGT A

31

10 (2) INFORMATION FOR SEQ ID NO: 5:

- (i) SEQUENCE CHARACTERISTICS:  
10 (A) LENGTH: 5 amino acids  
(B) TYPE: amino acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

15 (ii) MOLECULE TYPE: peptide

15 (iii) HYPOTHETICAL: YES

20 (iv) ANTI-SENSE: NO

25 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 5:

25 Leu Ile His Phe Gly  
1 5

30 (2) INFORMATION FOR SEQ ID NO: 6:

- 30 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 9 amino acids  
(B) TYPE: amino acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

35 (ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: YES

40 (iv) ANTI-SENSE: NO

45 (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 6:

45 Asp Tyr Glu Asp Arg Tyr Tyr Arg Glu  
1 5

50 (2) INFORMATION FOR SEQ ID NO: 7:

- 50 (i) SEQUENCE CHARACTERISTICS:  
(A) LENGTH: 7 amino acids  
(B) TYPE: amino acid  
(C) STRANDEDNESS: single  
(D) TOPOLOGY: linear

55 (ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: YES

(iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 7:

10       Gly Ser Asp Tyr Glu Asp Arg  
          1                   5

(2) INFORMATION FOR SEQ ID NO: 8:

15       (i) SEQUENCE CHARACTERISTICS:  
          (A) LENGTH: 9 amino acids  
          (B) TYPE: amino acid  
          (C) STRANDEDNESS: single  
          (D) TOPOLOGY: linear

20       (ii) MOLECULE TYPE: peptide

(iii) HYPOTHETICAL: YES

25       (iv) ANTI-SENSE: NO

30       (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 8:

      Tyr Tyr Arg Pro Val Asp Gln Tyr Ser  
          1                   5

35       (2) INFORMATION FOR SEQ ID NO: 9:

40       (i) SEQUENCE CHARACTERISTICS:  
          (A) LENGTH: 13 amino acids  
          (B) TYPE: amino acid  
          (C) STRANDEDNESS: single  
          (D) TOPOLOGY: linear

(ii) MOLECULE TYPE: peptide

45       (iii) HYPOTHETICAL: YES

(iv) ANTI-SENSE: NO

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(xi) SEQUENCE DESCRIPTION: SEQ ID NO: 9:

55       Cys Ile Thr Gln Tyr Gln Arg Glu Ser Gln Ala Tyr Tyr  
          1                   5                   10